

## SEQUENCE LISTING

Rotin, Daniela and Pham, Nam

```
<120> RAS Activator Nucleic Acid Molecules, Polypeptides and
      Methods of Use
<130> DDW-5001-US
<140> 09/911,826
<141> 2001-07-20
<150> PCT/CA00/00042
<151> 2000-01-20
<150> 2,259,830
<151> 1999-01-20
<160> 27
<170> PatentIn Ver. 2.1
<210> 1
<211> 6568
<212> DNA
<213> Homo sapiens
<220>
<221> CDS
<222> (63)..(4562)
<400> 1
cttgccatcg tgagagattg gtacatgatg tgtaaattca gttcagcata tgtttcttca 60
tt atg aaa cca cta gca atc cca gct aac cat gga gtt atg ggc cag
                                                                   107
   Met Lys Pro Leu Ala Ile Pro Ala Asn His Gly Val Met Gly Gln
cag gag aaa cac tca ctt cct gca gat ttc aca aaa ctg cat ctt act
                                                                   155
Gln Glu Lys His Ser Leu Pro Ala Asp Phe Thr Lys Leu His Leu Thr
                 20
gac agt ctc cac cca cag gtg acc cac gtt tct tct agc cat tca gga
                                                                   203
Asp Ser Leu His Pro Gln Val Thr His Val Ser Ser His Ser Gly
             35
tgt agt atc act agt gat tct ggg agc agc agt ctt tct gat atc tac
                                                                   251
Cys Ser Ile Thr Ser Asp Ser Gly Ser Ser Leu Ser Asp Ile Tyr
         50
cag gcc aca gaa agc gag gct ggt gat atg gac ctg aqt ggg ttg cca
                                                                   299
Gln Ala Thr Glu Ser Glu Ala Gly Asp Met Asp Leu Ser Gly Leu Pro
     65
                         70
gaa aca gca gtg gat tcc gaa gac gac gat gaa gaa gac att gag
Glu Thr Ala Val Asp Ser Glu Asp Asp Asp Glu Glu Asp Ile Glu
80
                     85
```

											aga Arg			395
											gaa Glu			443
											aca Thr 140			491
											gtg Val			539
											tcc Ser			587
						_				_	gga Gly		_	 635
											acc Thr			683
_		_				_	-		_	 _	gac Asp 220	_	_	731
_	_		_	_		_			_		aat Asn			779
_		_			_	_		-			gtt Val			827
_		_	_		_	_					gga Gly			875
											gtg Val			923
	_	_	_					_	_	_	ttg Leu 300			 971
											tta Leu			1019

					ctc Leu 325											1067	
					ttc Phe											1115	
_			_	_	ttt Phe	-				_				_		1163	
					ttg Leu											1211	
					aca Thr											1259	
					tct Ser 405		_						_	Āsp	_	1307	
					aaa Lys											1355	
					aat Asn											1403	
	-	_	_		ctt Leu	_										1451	
					gta Val											1499	
		_			gcc Ala 485								_			1547	
					tcc Ser											1595	
				_	aaa Lys											1643	
				_	ctg Leu											1691	
atc	ttg	cca	cag	aaa	cca	tac	aat	gat	att	999	att	ggt	cag	tct	caa	1739	

Ile	Leu 545	Pro	Gln	Lys	Pro	Tyr 550	Asn	Asp	Ile	Gly	Ile 555	Gly	Gln	Ser	Gln	
							agg Arg									1787
							tca Ser									1835
							ttc Phe									1883
							gct Ala 615									1931
							aag Lys									1979
							ccg Pro									2027
							atc Ile									2075
							ata Ile									2123
							act Thr 695									2171
							att Ile									2219
							atg Met									2267
							gat Asp									2315
							aga Arg									2363
							att Ile									2411

770 775 780

	atg Met 785	_			_				_		_	_		_		2459
_	tgc Cys	_							_							2507
_	gca Ala			_	_	_	_	_								2555
	tac Tyr															2603
_	aac Asn	_	-			_		_			_					2651
	ccc Pro 865							_			_	_				2699
	cac His															2747
_	cta Leu		_		_											2795
	gtg Val		_	-		_		_					_	_		2843
	cgg Arg	_	_				_	_		_			_			2891
	gat Asp 945															2939
	ttt Phe			-		-			_	_	_					2987
	gtg Val															3035
_	ctt Leu	_				_	_	_	Glu		-			Thr	_	3083

			aaa tcc gag ac Lys Ser Glu Th 1020	
		Gln Gln Lys	gct cag tcc ct Ala Gln Ser Le 1035	
	_	_	atc aac cag gg Ile Asn Gln Gl 1050	
	-		aag aaa gtg cc Lys Lys Val Pr 5	
_	o Phe Gly Ile		caa gct tta aa Gln Ala Leu Ly 10	s Lys Ile
_			cgt cac aag aa Arg His Lys Ly 1100	
	_	Ser Ser Gln	ctt tct tct cc Leu Ser Ser Pr 1115	
			act ttg gct cc Thr Leu Ala Pr 1130	
	_		agt gaa att tc Ser Glu Ile Se 5	<u> </u>
_	l Ser Asn Sei	_	tca gtg cca gt Ser Val Pro Va 11	l Ser Leu
		_	agc atc gtg ga Ser Ile Val Gl 1180	
cta ggg atg gg	agg atg gag	, agg cgg acc	atg att gaa cc	t gat cag 3659
1185	/ Arg Met Glu	Arg Arg Thr	Met Ile Glu Pr 1195	
1185 tat agc ttg gg	119 g tcc tat gca	Arg Arg Thr	Met Ile Glu Pr	c tta tat 3707

				Arg			ctt Leu		Ala					Arg		3803
			Ser				ggc Gly 1255	Ser					Gln			3851
		Gln					act Thr )					His				3899
	Tyr					Ala	ggt Gly				Ser					3947
					Ser		cat His			Lys					Asn	3995
				Ser			caa Gln		Gln					Trp		4043
			Gly				gaa Glu 1335	Asp					Thr			4091
		Arg					gat Asp )					Ala				4139
	Leu					Thr	gaa Glu				Pro					4187
					Ala		agt Ser			Lys					Arg	4235
				Tyr			ccc Pro		Pro					Tyr		4283
			Ile				cca Pro 1415	Glu					Pro			4331
	_	Pro	_				gcc Ala		_	_	_	Arg	_	_	_	4379
	Ser					Gly	cct Pro				Gln					4427
cat	ccc	acc	agc	agc	agg	cct	gtg	aac	aaa	cct	cag	tgg	cat	aaa	ccg	4475

His Pro Thr Ser Ser Arg Pro Val Asn Lys Pro Gln Trp His Lys Pro 1460 1465 1470

aac gag tot gac cog cgc otc gcc cot tat cag too caa ggg tot too 4523 Asn Glu Ser Asp Pro Arg Leu Ala Pro Tyr Gln Ser Gln Gly Phe Ser 1475 1480 1485

acc gag gag gat gaa gat gaa caa gtt tct gct gtt tga ggcacagact 4572 Thr Glu Glu Asp Glu Asp Glu Gln Val Ser Ala Val 1490 1495

tttctqqaaq caqaqcqaqc cacctqaaaq gagaqcacaa gaagacgtcc tgagcattgg 4632 agccttggaa ctcacattct gaggacggtg gaccagtttg cctccttccc tgccttaaaa 4692 gcagcatggg gcttcttctc cccttcttcc tttccccttt gcatgtgaaa tactgtgaag 4752 aaattgeeet ggeaetttte agaetttgtt gettgaaatg cacagtgeag caatettega 4812 gctcccactg ttgctgcctg ccacatcaca cagtatcatt ccaaattcca agatcatcac 4872 aacaagatga ttcactctgg ctgcacttct caatgcctgg aaggattttt tttaatcttc 4932 cttttagatt tcaatccagt cctagcactt gatctcattg ggataatgag aaaagctagc 4992 cattgaacta cttggggcct ttaacccacc aaggaagaca aagaaaaaca atgaaatcct 5052 ttgagtacag tgcttgtcca cttgtttaca atgtcctcct tttaaaaaaa aaaatgagtt 5112 taaagatttt gttcagagag taaatatata tccatttaat gattacagta ttattttaaa 5172 ccttaagtag ggttgccagc ctggtttctg aaaaaccaaa tatgccggac agggtgtggc 5232 cacaccaaga agacgggaag acctggcttg tgaccctggc ttcccatgtc cttctggtct 5292 caccegegaa gtgccctate etggaagtat gaaatgttag ccaattaata ccaagacace 5352 tcatctgctc cttccccagt ggatggggtt cttctgtaaa actgtttgca catggccagg 5412 ggagggaact aggaccettg tgteetgtet gageettatg gaggeaggae ggtgteattg 5472 gcggatgtgt cctgctccat tgagatggat ggcaaacccc atttttaagt tatatttctt 5532 tgatttttgt taatttagag gtgtaggttt tgttttttgt tttttgtttt tttttaagag 5592 aaacatttat aactggatag cattgcagtg aaagcagctt gggatgttgg agctaatgcc 5652 agctgtttat actgctcttt caagacagcc tccctttatt gaattggcat tagggaataa 5712 acaagcettt aaacgtgata aaagatcaaa aacctggtta gacatgccag cetttgcaag 5772 gcaggttagt caccaaagac taacctccaa gtggctttat ggacgctgca tatagagaag 5832 gcctaagtgt agcaaccatc tgctcacagc tgctattaac cctataatga ctgaaatgac 5892 ccctccactc tatttttgtg ttgttttgca cagactccgg aaaagtgaag gctgccaatc 5952

tgagtagtac tcaaatgtga ggaactgctg gtcttggatt tttttccat taaattcagc 6012
tgatcatatt gatcagtaga taaacgtaaa tagcttcaaa ttttaaaagt ggaattgcag 6072
tgtttttca ctgtatcaaa caatgtcagt gctttattta ataattctct tctgtatcat 6132
ggcatttgtc tacttgctta ttacattgtc aattatgcat ttgtaatttt acatgtaata 6192
tgcattattt gccagtttta ttatataggc tatggacctc atgtgcatat agaaagacag 6252
aaatctagct ctaccacaag ttgcacaaat gttatctaag cattaagtaa ttgtagaaca 6312
taggactgct aatctcagtt cgctctgtga tgtcaagtgc agaatgtaca attaactggt 6372
gatttcctca tactttgat actacttgta cctgtatgtc ttttagaaag acattggtgg 6432
agtctgtatc ccttttgtat ttttaataca ataattgtac atattggtta tattttgtt 6492
gaagatggta gaaatgtact atgtttatgc ttctacatcc agtttgtaca agctggaaaa 6552
taaataaata taacat

<210> 2

<211> 1499

<212> PRT

<213> Homo sapiens

<400> 2

Met Lys Pro Leu Ala Ile Pro Ala Asn His Gly Val Met Gly Gln Gln Glu Lys His Ser Leu Pro Ala Asp Phe Thr Lys Leu His Leu Thr Asp 25 Ser Leu His Pro Gln Val Thr His Val Ser Ser His Ser Gly Cys 40 Ser Ile Thr Ser Asp Ser Gly Ser Ser Leu Ser Asp Ile Tyr Gln 55 Ala Thr Glu Ser Glu Ala Gly Asp Met Asp Leu Ser Gly Leu Pro Glu 75 Thr Ala Val Asp Ser Glu Asp Asp Asp Glu Glu Asp Ile Glu Arg Ala Ser Asp Pro Leu Met Ser Arg Asp Ile Val Arg Asp Cys Leu Glu 105 Lys Asp Pro Ile Asp Arg Thr Asp Asp Ile Glu Gln Leu Leu Glu 120 Phe Met His Gln Leu Pro Ala Phe Ala Asn Met Thr Met Ser Val Arg 135 140 Arg Glu Leu Cys Ala Val Met Val Phe Ala Val Val Glu Arg Ala Gly 155 150 Thr Ile Val Leu Asn Asp Gly Glu Glu Leu Asp Ser Trp Ser Val Ile 170 Leu Asn Gly Ser Val Glu Val Thr Tyr Pro Asp Gly Lys Ala Glu Ile 185 Leu Cys Met Gly Asn Ser Phe Gly Val Ser Pro Thr Met Asp Lys Glu Tyr Met Lys Gly Val Met Arg Thr Lys Val Asp Asp Cys Gln Phe Val 215

Cys Ile Ala Gln Gln Asp Tyr Cys Arg Ile Leu Asn Gln Val Glu Lys Asn Met Gln Lys Val Glu Glu Glu Glu Ile Val Met Val Lys Glu His Arg Glu Leu Asp Arg Thr Gly Thr Arg Lys Gly His Ile Val Ile Lys Gly Thr Ser Glu Arg Leu Thr Met His Leu Val Glu Glu His Ser Val Val Asp Pro Thr Phe Ile Glu Asp Phe Leu Leu Thr Tyr Arg Thr Phe Leu Ser Ser Pro Met Glu Val Gly Lys Lys Leu Leu Glu Trp Phe Asn Asp Pro Ser Leu Arg Asp Lys Val Thr Arg Val Val Leu Leu Trp Val Asn Asn His Phe Asn Asp Phe Glu Gly Asp Pro Ala Met Thr Arg Phe Leu Glu Glu Phe Glu Asn Asn Leu Glu Arg Glu Lys Met Gly Gly His Leu Arg Leu Leu Asn Ile Ala Cys Ala Ala Lys Ala Lys Arg Arg Leu Met Thr Leu Thr Lys Pro Ser Arg Glu Ala Pro Leu Pro Phe Ile Leu Leu Gly Gly Ser Glu Lys Gly Phe Gly Ile Phe Val Asp Ser Val Asp Ser Gly Ser Lys Ala Thr Glu Ala Gly Leu Lys Arg Gly Asp Gln Ile Leu Glu Val Asn Gly Gln Asn Phe Glu Asn Ile Gln Leu Ser Lys Ala Met Glu Ile Leu Arg Asn Asn Thr His Leu Ser Ile Thr Val Lys Thr Asn Leu Phe Val Phe Lys Glu Leu Leu Thr Arg Leu Ser Glu Glu Lys Arg Asn Gly Ala Pro His Leu Pro Lys Ile Gly Asp Ile Lys Lys Ala Ser Arg Tyr Ser Ile Pro Asp Leu Ala Val Asp Val Glu Gln Val Ile Gly Leu Glu Lys Val Asn Lys Lys Ser Lys Ala Asn Thr Val Gly Gly Arg Asn Lys Leu Lys Lys Ile Leu Asp Lys Thr Arg Ile Ser Ile Leu Pro Gln Lys Pro Tyr Asn Asp Ile Gly Ile Gly Gln Ser Gln Asp Asp Ser Ile Val Gly Leu Arg Gln Thr Lys His Ile Pro Thr Ala Leu Pro Val Ser Gly Thr Leu Ser Ser Ser Asn Pro Asp Leu Leu Gln Ser His His Arg Ile Leu Asp Phe Ser Ala Thr Pro Asp Leu Pro Asp Gln Val Leu Arg Val Phe Lys Ala Asp Gln Gln Ser Arg Tyr Ile Met Ile Ser Lys Asp Thr Thr Ala Lys Glu Val Val Ile Gln Ala Ile Arg Glu Phe Ala Val Thr Ala Thr Pro Asp Gln Tyr Ser Leu Cys Glu Val Ser Val Thr Pro Glu Gly Val Ile Lys Gln Arg Arg Leu Pro Asp Gln Leu Ser Lys Leu Ala Asp Arg Ile Gln Leu Ser Gly Arg Tyr Tyr Leu Lys

	675					680					685			
Asn Asn		Glu	Thr	Glu	Thr		Cys	Ser	Asp	Glu		Ala	Gln	Glu
690					695					700				
Leu Leu	Arg	Glu	Ser		Ile	Ser	Leu	Leu		Leu	Ser	Thr	Val	
705	1	~ 7	_	710		_	_	-1	715	_	-1	_	•	720
Val Ala	Thr	GIn		Ser	Met	Arg	Asn		GIu	Leu	Pne	Arg		11e
Glu Pro	Thr	Glu	725	Tle	Δen	Δen	T.e.11	730 Dhe	Lare	Len	Δrα	Ser	735	Thr
GIU FIO	TIIL	740	TYT	110	тор	ASP	745	1110	цуз	пси	nr 9	750	цуБ	1111
Ser Cys	Ala 755		Leu	Lys	Arg	Phe 760		Glu	Val	Ile	Asn 765		Glu	Thr
Phe Trp		Ala	Ser	Glu	Ile 775		Arg	Glu	Thr	Asn 780		Leu	Lys	Arg
Met Lys	Ile	Ile	Lys	His		Ile	Lys	Ile	Ala		His	Cys	Arg	Glu
785				790					795					800
Cys Lys	Asn	Phe	Asn 805	Ser	Met	Phe	Ala	Ile 810	Ile	Ser	Gly	Leu	Asn 815	Leu
Ala Pro	Val	Ala 820	Arg	Leu	Arg	Thr	Thr 825	Trp	Glu	Lys	Leu	Pro 830	Asn	Lys
Tyr Glu	Lys 835		Phe	Gln	Asp	Leu 840		Asp	Leu	Phe	Asp 845	Pro	Ser	Arg
Asn Met		Lys	Tyr	Arg			Leu	Asn	Ser	Gln 860		Leu	Gln	Pro
850 Pro Ile	Tle	Pro	Len	Phe	855 Pro	Val	Tle	Lvs	Lvs		Len	Thr	Phe	Len
865	110	110	пси	870	110	vai	110	my b	875	тор	шец	1111	1110	880
His Glu	Gly	Asn	Asp 885	Ser	Lys	Val	Asp	Gly 890	Leu	Val	Asn	Phe	Glu 895	Lys
Leu Arg	Met	Ile 900		Lys	Glu	Ile	Arg 905		Val	Gly	Arg	Met 910		Ser
Val Asn	Met 915		Pro	Ala	Leu	Met 920		Arg	Thr	Arg	Lys 925		Lys	Trp
Arg Ser 930		Gly	Ser	Leu	Ser 935		Gly	Ser	Thr	Asn 940		Thr	Val	Leu
Asp Val	Ala	Gln	Thr	Gly		His	Lys	Lys	Arg	-	Arg	Arg	Ser	Ser
945		_		950			_		955	_		_		960
Phe Leu	Asn	Ala	1965 965	Lys	Leu	Tyr	Glu	Asp 970	Ala	Gln	Met	Ala	Arg 975	Lys
Val Lys	Gln	_	Leu	Ser	Asn	Leu		Leu	Glu	Met	Asp		Glu	Ser
Leu Gln	Thr	980	Sar	T. 211	Gln	Cvc	985	Dro	Λla	Thr	λan	990	T.011	Dro
neu Gin	995	пеи	SEI	пеп		L000	Giu	PIO	Ата		1005	1111	Бец	PIO
Lys Asn		Gly	Asp	Lys			Val	Lys	Ser			Ser	Pro	Val
1010				1	.015				1	1020				
Ala Pro	Arg	Ala	_		Gln	Gln	Lys			Ser	Leu	Pro		
1025	<b>a</b> 1	D		L030	<b>3</b> .7 -	77.	T		1035	~1	<b>a</b> 1	T		1040
Gln Gln		1	045				_ 1	1050				1	.055	
Pro Ala		Ser 1060	Leu	Tyr	Pro		Arg 1065	Lys	Lys	Val		Val L070	Lys	Asp
Leu Pro	Pro 1075	Phe	Gly	Ile		Ser 1080	Pro	Gln	Ala		Lys .085	Lys	Ile	Leu
Ser Leu		Glu	Glu	_	Ser		Glu	Arg		Lys		Gln	Ala	Glu
1090 Asp Thr	Tle	Ser	Asn		.095 Ser	Ser	Gln	Leu		.100 Ser	Pro	Pro	Thr	Ser
1105		501		110	J01	J-01			.115	JUL	110	110		.120
Pro Gln	Ser	Ser			Lys	Gly		Thr		Ala	Pro	Ser		
		1	.125				1	130				1	135	

Val Asp Asn Phe Ser Asp Ser Gly His Ser Glu Ile Ser Ser Arg Ser 1145 Ser Ile Val Ser Asn Ser Ser Phe Asp Ser Val Pro Val Ser Leu His 1160 1165 Asp Glu Arg Arg Gln Arg His Ser Val Ser Ile Val Glu Thr Asn Leu 1175 1180 1170 Gly Met Gly Arg Met Glu Arg Arg Thr Met Ile Glu Pro Asp Gln Tyr 1190 1195 Ser Leu Gly Ser Tyr Ala Pro Met Ser Glu Gly Arg Gly Leu Tyr Ala 1210 1205 Thr Ala Thr Val Ile Ser Ser Pro Ser Thr Glu Glu Leu Ser Gln Asp 1220 1225 Gln Gly Asp Arg Ala Ser Leu Asp Ala Ala Asp Ser Gly Arg Gly Ser 1240 1245 Trp Thr Ser Cys Ser Ser Gly Ser His Asp Asn Ile Gln Thr Ile Gln 1255 1260 His Gln Arg Ser Trp Glu Thr Leu Pro Phe Gly His Thr His Phe Asp 1270 1275 Tyr Ser Gly Asp Pro Ala Gly Leu Trp Ala Ser Ser His Met Asp 1290 1295 1285 Gln Ile Met Phe Ser Asp His Ser Thr Lys Tyr Asn Arg Gln Asn Gln 1305 1310 Ser Arg Glu Ser Leu Glu Gln Ala Gln Ser Arg Ala Ser Trp Ala Ser 1320 1325 1315 Ser Thr Gly Tyr Trp Gly Glu Asp Ser Glu Gly Asp Thr Gly Thr Ile 1335 Lys Arg Arg Gly Gly Lys Asp Val Ser Ile Glu Ala Glu Ser Ser Ser 1350 1355 Leu Thr Ser Val Thr Thr Glu Glu Thr Lys Pro Val Pro Met Pro Ala 1370 1365 His Ile Ala Val Ala Ser Ser Thr Thr Lys Gly Leu Ile Ala Arg Lys 1380 1385 1390 Glu Gly Arg Tyr Arg Glu Pro Pro Pro Thr Pro Pro Gly Tyr Ile Gly 1395 1400 1405 Ile Pro Ile Thr Asp Phe Pro Glu Gly His Ser His Pro Ala Arg Lys 1415 1420 Pro Pro Asp Tyr Asn Val Ala Leu Gln Arg Ser Arg Met Val Ala Arg 1430 1435 Ser Ser Asp Thr Ala Gly Pro Ser Ser Val Gln Gln Pro His Gly His 1445 1450 Pro Thr Ser Ser Arg Pro Val Asn Lys Pro Gln Trp His Lys Pro Asn 1465 1470 Glu Ser Asp Pro Arg Leu Ala Pro Tyr Gln Ser Gln Gly Phe Ser Thr 1480 Glu Glu Asp Glu Asp Glu Gln Val Ser Ala Val

<210> 3 <211> 799 <212> DNA

<213> Mus musculus

<400> 3

actaaaggga acaaaagctg gagctccacc gcggtggcgg ccgctctaga actagtggat 60 cccccgggct gcaggaattc aagcggtggg aaggatgtct ccgctgaggc agagagcagc 120

agcatggtgc	ccgtgactac	agaggaagcc	aaacctgtcc	ctatgcctgc	ccacatagct	180
gtgacgccga	gcactaccaa	gggactcatc	gcacggaagg	aaggcaggta	ccgggagccg	240
cctcccacac	ctccaggcta	cgtgggcatc	cccattgccg	atttcccaga	agggccttgc	300
cacccggcca	ggaagccccc	ggattacaac	gtggccctgc	agcggtcccg	catggtggca	360
cggcccactg	aggccccggc	accgggccag	acgccgcctg	cagccgcagc	cagccggccg	420
ggcagcaagc	cacagtggca	caagcccagc	gacgcagacc	cacgcctcgc	gcccttccag	480
gcaggcttcg	caggagcgga	ggaggacgaa	gatgaacaag	tgtctgctgt	ttgaggcgca	540
ggctccttga	tccacagtga	gccacccaaa	ggagagcaca	agaagacgtc	ccaagccttg	600
gagccttggc	acgcacatct	gaggatggtg	gaccagtttg	cctccttccc	tgccttaaag	660
cagcatgggg	cttcttctcc	ccttcttcct	ttcccctttg	catgtgaaat	actgtgaaga	720
aattgccctg	gcactttgca	gacttgttgc	ttgaaatgca	cagcccagca	gcccctgagc	780
tgctgcctgc	cacgtcacg					799

<210> 4

<211> 286

<212> PRT

<213> Homo sapiens

<220>

<221> SIMILAR

<222>

<223> Xaa is any aa

<400> 4

Thr Lys Gly Asn Lys Ser Trp Ser Ser Thr Ala Val Ala Ala Ala Leu 1 5 10 15

Glu Leu Val Asp Pro Pro Gly Cys Arg Asn Ser Ser Gly Gly Lys Asp
20 25 30

Val Ser Ala Glu Ala Glu Ser Ser Ser Met Val Pro Val Thr Thr Glu 35 40 45

Glu Ala Lys Pro Val Pro Met Pro Ala His Ile Ala Val Thr Pro Ser 50 55 60

Thr Thr Lys Gly Leu Ile Ala Arg Lys Glu Gly Arg Tyr Arg Glu Pro 65 70 75 80

Pro Pro Thr Pro Pro Gly Tyr Val Gly Ile Pro Ile Ala Asp Phe Pro 85 90 95

Glu Gly Pro Cys His Pro Ala Arg Lys Pro Pro Asp Tyr Asn Val Ala 100 105 110

Leu Gln Arg Ser Arg Met Val Ala Arg Pro Thr Glu Ala Pro Ala Pro 115 120 125

Gly Gln Thr Pro Pro Ala Ala Ala Ser Arg Pro Gly Ser Lys Pro 130 135 140

Gln Trp His Lys Pro Ser Asp Ala Asp Pro Arg Leu Ala Pro Phe Gln 145 150 155 160

Ala Ala Ser His Ser Gly Thr Ser Pro Ala Thr Gln Thr His Ala Ser 165 170 175 Arg Pro Ser Arg Gln Ala Ser Gln Glu Arg Arg Arg Thr Lys Met Asn 180 185 190

Lys Cys Leu Leu Phe Glu Ala Gln Ala Pro Xaa Ser Thr Val Ser His 195 200 205

Pro Lys Glu Ser Thr Arg Arg Arg Pro Lys Pro Trp Ser Leu Gly Thr 210 215 220

His Ile Xaa Gly Trp Trp Thr Ser Leu Pro Pro Ser Leu Pro Xaa Ser 225 230 235 240

Ser Met Gly Leu Leu Pro Phe Phe Leu Ser Pro Leu His Val Lys 245 250 255

Tyr Cys Glu Glu Ile Ala Leu Ala Leu Cys Arg Leu Val Ala Xaa Asn 260 265 270

Ala Gln Pro Ser Ser Pro Xaa Ala Ala Ala Cys His Val Thr 275 280 285

<210> 5

<211> 245

<212> PRT

<213> Homo sapiens

<220>

<221> SIMILAR

<222>

<223> Xaa is any aa

<400> 5

Leu Lys Gly Thr Lys Ala Gly Ala Pro Pro Arg Trp Arg Pro Leu Xaa 1 5 10 15

Asn Xaa Trp Ile Pro Arg Ala Ala Gly Ile Gln Ala Val Gly Arg Met 20 25 30

Ser Pro Leu Arg Gln Arg Ala Ala Trp Cys Pro Xaa Leu Gln Arg 35 40 45

Lys Pro Asn Leu Ser Leu Cys Leu Pro Thr Xaa Leu Xaa Arg Arg Ala 50 55 60

Leu Pro Arg Asp Ser Ser His Gly Arg Lys Ala Gly Thr Gly Ser Arg 65 70 75 80

Leu Pro His Leu Gln Ala Thr Trp Ala Ser Pro Leu Pro Ile Ser Gln 85 90 95

Lys Gly Leu Ala Thr Arg Pro Gly Ser Pro Arg Ile Thr Thr Trp Pro
100 105 110

Cys Ser Gly Pro Ala Trp Trp His Gly Pro Leu Arg Pro Arg His Arg 115 120 125 Ala Arg Arg Arg Leu Gln Pro Gln Pro Ala Gly Arg Arg Leu Arg Arg 130 135 140

Ser Gly Gly Gly Arg Arg Xaa Thr Ser Val Cys Cys Leu Arg Arg 145 150 155 160

Leu Leu Asp Pro Gln Xaa Ala Thr Gln Arg Arg Ala Gln Glu Asp Val 165 170 175

Pro Ser Leu Gly Ala Leu Ala Arg Thr Ser Glu Asp Gly Gly Pro Val 180 185 190

Cys Leu Leu Pro Cys Leu Lys Ala Ala Trp Gly Phe Phe Ser Pro Ser 195 200 205

Ser Phe Pro Leu Cys Met Xaa Asn Thr Val Lys Lys Leu Pro Trp His 210 215 220

Phe Ala Asp Leu Leu Glu Met His Ser Pro Ala Ala Pro Glu Leu 225 230 235 240

Leu Pro Ala Thr Ser 245

<210> 6

<211> 266

<212> PRT

<213> Homo sapiens

<220>

<221> SIMILAR

<222>

<223> Xaa is any aa

<400> 6

Xaa Arg Glu Gln Lys Leu Glu Leu His Arg Gly Gly Arg Ser Arg
1 5 10 15

Thr Ser Gly Ser Pro Gly Leu Gln Glu Phe Lys Arg Trp Glu Gly Cys
20 25 30

Leu Arg Xaa Gly Arg Glu Gln Gln His Gly Ala Arg Asp Tyr Arg Gly
35 40 45

Ser Gln Thr Cys Pro Tyr Ala Cys Pro His Ser Cys Asp Ala Glu His 50 55 60

Tyr Gln Gly Thr His Arg Thr Glu Gly Arg Gln Val Pro Gly Ala Ala 65 70 75 80

Ser His Thr Ser Arg Leu Arg Gly His Pro His Cys Arg Phe Pro Arg

Arg Ala Leu Pro Pro Gly Gln Glu Ala Pro Gly Leu Gln Arg Gly Pro
100 105 110

\_\_\_\_

Ala Ala Val Pro His Gly Gly Thr Ala His Xaa Gly Pro Gly Thr Gly 115 120 125

Pro Asp Ala Ala Cys Ser Arg Ser Gln Pro Ala Gly Gln Gln Ala Thr 130 135 140

Val Ala Gln Ala Gln Arg Arg Pro Thr Pro Arg Ala Leu Pro Gly
145 150 155 160

Ala Gly Phe Ala Gly Ala Glu Glu Asp Glu Asp Glu Gln Val Ser Ala 165 170 175

Val Xaa Gly Ala Gly Ser Leu Ile His Ser Glu Pro Pro Lys Gly Glu 180 185 190

His Lys Lys Thr Ser Gln Ala Leu Glu Pro Trp His Ala His Leu Arg 195 200 205

Met Val Asp Gln Phe Ala Ser Phe Pro Ala Leu Lys Gln His Gly Ala 210 215 220

Ser Ser Pro Leu Leu Pro Phe Pro Phe Ala Cys Glu Ile Leu Xaa Arg 225 230 235 240

Asn Cys Pro Gly Thr Leu Gln Thr Cys Cys Leu Lys Cys Thr Ala Gln 245 250 255

Gln Pro Leu Ser Cys Cys Leu Pro Arg His 260 265

<210> 7

<211> 307

<212> PRT

<213> Drosophila melanogaster

<400> 7

Ser Asn Val His Phe Leu His Leu Asn Ala Tyr Glu Leu Ala Ile Gln
1 5 10 15

Leu Thr Leu Gln Asp Phe Ala Asn Phe Arg Gln Ile Glu Ser Thr Glu
20 25 30

Tyr Val Asp Glu Leu Phe Glu Leu Arg Ser Arg Tyr Gly Val Pro Met
35 40 45

Leu Ser Lys Phe Ala Glu Leu Val Asn Arg Glu Met Phe Trp Val Val
50 60

Ser Glu Ile Cys Ala Glu His Asn Ile Val Arg Arg Met Lys Ile Val 65 70 75 80

Lys Gln Phe Ile Lys Ile Ala Arg His Cys Lys Glu Cys Arg Asn Phe 85 90 95

Asn Ser Met Phe Ala Ile Val Ser Gly Leu Gly His Gly Ala Val Ser

100 105 110

Arg Leu Arg Gln Thr Trp Glu Lys Leu Pro Ser Lys Tyr Gln Arg Leu 115 120 125

Phe Asn Asp Leu Gln Asp Leu Met Asp Pro Ser Arg Asn Met Ser Lys 130 135 140

Tyr Arg Gln Leu Val Ser Ala Glu Leu Leu Ala Gln His Pro Ile Ile 145 150 155 160

Pro Phe Tyr Pro Ile Val Lys Lys Asp Leu Thr Phe Ile His Leu Gly
165 170 175

Asn Asp Thr Arg Val Asp Gly Leu Val Asn Phe Glu Lys Leu Arg Met 180 185 190

Leu Ala Lys Glu Val Arg Leu Leu Thr His Met Cys Ser Ser Pro Tyr
195 200 205

Asp Leu Leu Ser Ile Leu Glu Leu Lys Gly Gln Ser Pro Ser Asn Ala 210 215 220

Leu Phe Ser Leu Asn Gln Met Ser Ala Ser Gln Ser Asn Ala Ala Ala 225 230 235 240

Gly Thr Val Ile Ala Ala Asn Ala Gly Gln Ala Thr Ile Lys Arg Arg 245 250 255

Lys Lys Ser Thr Ala Ala Pro Asn Pro Lys Lys Met Phe Glu Glu Ala 260 265 270

Gln Met Val Arg Arg Val Lys Ala Tyr Leu Asn Ser Leu Lys Ile Leu 275 280 285

Ser Asp Glu Asp Leu Leu His Lys Phe Ser Leu Glu Cys Glu Pro Ala 290 295 300

His Gly Ser

<210> 8

<211> 270

<212> PRT

<213> Homo sapiens

<400> 8

Ser Ala Glu Gly Leu Asp Leu Val Ser Ala Lys Asp Leu Ala Gly Gln
1 5 10 15

Leu Thr Asp His Asp Trp Ser Leu Phe Asn Ser Ile His Gln Val Glu 20 25 30

Leu Ile His Tyr Val Leu Gly Pro Gln His Leu Arg Asp Val Thr Thr 35 40 45

Ala Asn Leu Glu Arg Phe Met Arg Arg Phe Asn Glu Leu Gln Tyr Trp 55 Val Ala Thr Glu Leu Cys Leu Cys Pro Val Pro Gly Pro Arg Ala Gln 70 Leu Leu Arg Lys Phe Ile Lys Leu Ala Ala His Leu Lys Glu Gln Lys Asn Leu Asn Ser Phe Phe Ala Val Met Phe Gly Leu Ser Asn Ser Ala 105 Ile Ser Arg Leu Ala His Thr Trp Glu Arg Leu Pro His Lys Val Arg 120 Lys Leu Tyr Ser Ala Leu Glu Arg Leu Leu Asp Pro Ser Trp Asn His 135 Arg Val Tyr Arg Leu Ala Leu Ala Lys Leu Ser Pro Pro Val Ile Pro 150 Phe Met Pro Leu Leu Lys Asp Met Thr Phe Ile His Glu Gly Asn His Thr Leu Val Glu Asn Leu Ile Asn Phe Glu Lys Met Arg Met Met 185 Ala Arg Ala Ala Arg Met Leu His His Cys Arg Ser His Asn Pro Val 200 Pro Leu Ser Pro Leu Arg Ser Arg Val Ser His Leu His Glu Asp Ser 215 220 Gln Val Ala Arg Ile Ser Thr Cys Ser Glu Gln Ser Leu Ser Thr Arg 235 230 Ser Pro Ala Ser Thr Trp Ala Tyr Val Gln Gln Leu Lys Val Ile Asp 250 245 Asn Gln Arg Glu Leu Ser Arg Leu Ser Arg Glu Leu Glu Pro 260 265

<210> 9

<211> 244

<212> PRT

<213> Mus musculus

<400> 9

Lys Ala Glu Cys Phe Glu Thr Leu Ser Ala Met Glu Leu Ala Glu Gln 1 5 10 15

Ile Thr Leu Leu Asp His Ile Val Phe Arg Ser Ile Pro Tyr Glu Glu
20 25 30

Phe Leu Gly Gln Gly Trp Met Lys Leu Asp Lys Asn Glu Arg Thr Pro

Tyr Ile Met Lys Thr Ser Gln His Phe Asn Glu Met Ser Asn Leu Val 50 55 60

Ala Ser Gln Ile Met Asn Tyr Ala Asp Ile Ser Ser Arg Pro Asn Ala 65 70 75 80

Ile Glu Lys Trp Val Ala Val Ala Asp Ile Cys Arg Cys Leu His Asn 85 90 95

Tyr Asn Gly Val Leu Glu Ile Thr Ser Ala Leu Asn Arg Ser Pro Ile 100 105 110

Tyr Arg Leu Lys Lys Thr Trp Ala Lys Val Ser Lys Gln Thr Lys Ala 115 120 125

Leu Met Asp Lys Leu Gln Lys Thr Val Ser Ser Glu Gly Arg Phe Lys 130 135 140

Asn Leu Arg Glu Thr Leu Lys Asn Cys Asn Pro Pro Ala Val Pro Tyr 145 150 155 160

Leu Gly Met Tyr Leu Thr Asp Leu Ala Phe Ile Glu Glu Gly Thr Pro 165 170 175

Asn Phe Thr Glu Glu Gly Leu Val Asn Phe Ser Lys Met Arg Met Ile 180 185 190

Ser His Ile Ile Arg Glu Ile Arg Gln Phe Gln Gln Thr Ala Tyr Arg 195 200 205

Ile Asp Gln Gln Pro Lys Val Ile Gln Tyr Leu Leu Asp Lys Ala Leu 210 215 220

Val Ile Asp Glu Asp Ser Leu Tyr Glu Leu Ser Leu Lys Ile Glu Pro 225 230 235 240

Arg Leu Pro Ala

<210> 10

<211> 249

<212> PRT

<213> Homo sapiens

<400> 10

Asp Glu Ile Thr Leu Leu Thr Leu His Pro Leu Glu Leu Ala Arg Gln
1 5 10 15

Leu Thr Leu Leu Glu Phe Glu Met Tyr Lys Asn Val Lys Pro Ser Glu 20 25 30

Leu Val Gly Ser Pro Trp Thr Lys Lys Asp Lys Glu Val Lys Ser Pro

Asn Leu Leu Lys Ile Met Lys His Thr Thr Asn Val Thr Arg Trp Ile

50 55 60

Glu Lys Ser Ile Thr Glu Ala Glu Asn Tyr Glu Glu Arg Leu Ala Ile 70 Met Gln Arg Ala Ile Glu Val Met Met Val Met Leu Glu Leu Asn Asn 90 Phe Asn Gly Ile Leu Ser Ile Val Ala Ala Met Gly Thr Ala Ser Val 100 Tyr Arg Leu Arg Trp Thr Phe Gln Gly Leu Pro Glu Arg Tyr Arg Lys Phe Leu Glu Glu Cys Arg Glu Leu Ser Asp Asp His Leu Lys Lys Tyr 135 Gln Glu Arg Leu Arg Ser Ile Asn Pro Pro Cys Val Pro Phe Phe Gly 155 150 Arg Tyr Leu Thr Asn Ile Leu His Leu Glu Glu Gly Asn Pro Asp Leu 165 170 Leu Ala Asn Thr Glu Leu Ile Asn Phe Ser Lys Arg Arg Lys Val Ala 180 185 Glu Ile Ile Gly Glu Ile Gln Gln Tyr Gln Asn Gln Pro Tyr Cys Leu 200 Asn Glu Glu Ser Thr Ile Arg Gln Phe Phe Glu Gln Leu Asp Pro Phe 215 Asn Gly Leu Ser Asp Lys Gln Met Ser Asp Tyr Leu Tyr Asn Glu Ser 225 230 235 Leu Arg Ile Glu Pro Arg Gly Cys Lys 245 <210> 11 <211> 243 <212> PRT <213> Homo sapiens <400> 11

Val Ser Leu Leu Phe Asp His Leu Glu Pro Glu Glu Leu Ser Glu His
1 5 10 15

Leu Thr Tyr Leu Glu Phe Lys Ser Phe Arg Arg Ile Ser Phe Ser Asp
20 25 30

Tyr Gln Asn Tyr Leu Val Asn Ser Cys Val Lys Glu Asn Pro Thr Met
35 40 45

Glu Arg Ser Ile Ala Leu Cys Asn Gly Ile Ser Gln Trp Val Gln Leu 50 55 60

Met Val Leu Ser Arg Pro Thr Pro Gln Leu Arg Ala Glu Val Phe Ile Lys Phe Ile Gln Val Ala Gln Lys Leu His Gln Leu Gln Asn Phe Asn 90 Thr Leu Met Ala Val Ile Gly Gly Leu Cys His Ser Ser Ile Ser Arg 105 100 Leu Lys Glu Thr Ser Ser His Val Pro His Glu Ile Asn Lys Val Leu 120 Gly Glu Met Thr Glu Leu Leu Ser Ser Ser Arg Asn Tyr Asp Asn Tyr Arg Arg Ala Tyr Gly Glu Cys Thr Asp Phe Lys Ile Pro Ile Leu Gly 150 Val His Leu Lys Asp Leu Ile Ser Leu Tyr Glu Ala Met Pro Asp Tyr 170 Leu Glu Asp Gly Lys Val Asn Val His Lys Leu Leu Ala Leu Tyr Asn 180 185 His Ile Ser Glu Leu Val Gln Leu Gln Glu Val Ala Pro Pro Leu Glu 200 Ala Asn Lys Asp Leu Val His Leu Leu Thr Leu Ser Leu Asp Leu Tyr 215 220 Tyr Thr Glu Asp Glu Ile Tyr Glu Leu Ser Tyr Ala Arg Glu Pro Arg 225 230 235 Asn His Arg <210> 12 <211> 48 <212> PRT <213> Unknown Organism <220> <223> Description of Unknown Organism: unavailable <400> 12 Ile Arg Gly Gly Thr Lys Glu Ala Leu Ile Glu His Leu Thr Ser His Glu Leu Val Asp Ala Ala Phe Asn Val Thr Met Leu Ile Thr Phe Arg 20

Ser Ile Leu Thr Thr Arg Glu Phe Phe Tyr Ala Leu Ile Tyr Arg Tyr

45

40

```
<211> 47
<212> PRT
<213> Mus musculus
<400> 13
Ile Lys Gly Gly Thr Val Val Lys Leu Ile Glu Arg Leu Thr Tyr His
                  5
Met Tyr Ala Asp Pro Asn Phe Val Arg Thr Phe Leu Thr Tyr Arg Ser
Phe Cys Lys Gln Glu Leu Leu Asn Leu Leu Ile Glu Arg Phe Glu
<210> 14
<211> 48
<212> PRT
<213> Mus musculus
<400> 14
Ile Arg Tyr Ala Ser Val Glu Ala Leu Leu Glu Arg Leu Thr Asp Leu
Arg Phe Leu Ser Ile Asp Phe Leu Asn Thr Phe Leu His Thr Tyr Arg
                                 25
Ile Phe Thr Thr Ala Thr Val Val Leu Ala Lys Leu Ser Asp Ile Tyr
                             40
         35
<210> 15
<211> 50
<212> PRT
<213> Unknown Organism
<220>
<221> SIMILAR
<222>
<223> Xaa is any aa
<220>
<223> Description of Unknown Organism: unavailable
```

20

5

<400> 15

<210> 13

10

Val Val Lys Phe Ala Ser Leu Asn Lys Leu Val Glu His Leu Thr His

Asp Ser Lys His Asp Leu Gln Phe Leu Lys Thr Phe Leu Met Thr Tyr

25

Gln Ser Phe Cys Thr Pro Glu Lys Leu Met Ser Lys Leu Gln Gln Arg 40 Tyr Xaa 50 <210> 16 <211> 77 <212> PRT <213> Drosophila melanogaster <400> 16 Leu Thr Arg Ser Ser Arg Asp Glu Pro Leu Asn Phe Arg Ile Val Gly 10 Gly Tyr Glu Leu Arg Gly Val Ala Ile Ala Thr Gly Asn Ala Ala Val Gly Ile Tyr Ile Ser His Val Glu Pro Gly Ser Lys Ala Gln Asp Val Gly Leu Lys Arg Gly Asp Gln Ile His Glu Val Asn Gly Gln Ser Leu Asp His Val Thr Ser Lys Arg Ala Leu Glu Ile Leu Thr 70 <210> 17 <211> 71 <212> PRT <213> Homo sapiens <400> 17 Asn Leu Lys Lys Asp Ala Lys Tyr Gly Leu Gly Phe Gln Ile Ile Gly Gly Glu Lys Met Gly Arg Leu Asp Leu Gly Ile Phe Ile Ser Ser Val 20 Ala Pro Gly Gly Pro Ala Asp Leu Asp Gly Cys Leu Lys Pro Gly Asp 35 Arg Leu Ile Ser Val Asn Ser Val Ser Leu Glu Gly Val Ser His His Ala Ala Ile Glu Ile Leu Gln 65 70 <210> 18 <211> 67 <212> PRT <213> Homo sapiens <400> 18

Ile Val Ile His Arg Gly Ser Thr Gly Leu Gly Phe Asn Ile Val Gly Gly Glu Asp Gly Glu Gly Ile Phe Ile Ser Phe Ile Leu Ala Gly Gly 25 Pro Ala Asp Leu Ser Gly Glu Leu Arg Lys Gly Asp Gln Ile Leu Ser Val Asn Gly Val Asp Leu Arg Asn Ala Ser His Glu Gln Ala Ala Ile 55 Ala Leu Lys 65 <210> 19 <211> 68 <212> PRT <213> Rattus rattus <400> 19 Val Glu Leu Pro Lys Thr Glu Glu Gly Leu Gly Phe Asn Ile Met Gly Gly Lys Glu Gln Asn Ser Pro Ile Tyr Ile Ser Arg Ile Ile Pro Gly Gly Ile Ala Asp Arg His Gly Gly Leu Lys Arg Gly Asp Gln Leu Leu 35 40 Ser Val Asn Gly Val Ser Val Glu Gly Glu His His Glu Lys Ala Val Glu Leu Leu Lys 65 <210> 20 <211> 65 <212> PRT <213> Homo sapiens <400> 20 Val Lys Val Gln Lys Gly Ser Glu Pro Leu Gly Ile Ser Ile Val Ser Gly Glu Lys Gly Gly Ile Tyr Val Ser Lys Val Thr Val Gly Ser Ile Ala His Gln Ala Gly Leu Glu Tyr Gly Asp Gln Leu Leu Glu Phe Asn 40 Gly Ile Asn Leu Arg Ser Ala Thr Glu Gln Ala Arg Leu Ile Ile 55

Gly

<210> 21

<211> 98

<212> PRT

<213> Drosophila melanogaster

<400> 21

Met Val Phe Ala Val Val Asp Lys Ala Gly Thr Val Val Met Ser Asp 1 5 10 15

Gly Glu Glu Leu Asp Ser Trp Ser Val Leu Ile Asn Gly Ala Val Glu 20 25 30

Ile Glu His Ala Asn Gly Ser Arg Glu Glu Leu Gln Met Gly Asp Ser 35 40 45

Phe Gly Ile Leu Pro Thr Met Asp Lys Leu Tyr His Arg Gly Val Met 50 55 60

Arg Thr Lys Cys Asp Asp Cys Gln Phe Val Cys Ile Thr Gln Thr Asp 65 70 75 80

Tyr Tyr Arg Ile Gln His Gln Gly Glu Glu Asn Thr Arg Arg His Glu
85 90 95

Asp Glu

<210> 22

<211> 99

<212> PRT

<213> Homo sapiens

<400> 22

Leu Leu Phe Glu Pro His Ser Lys Ala Gly Thr Val Leu Phe Ser Gln 1 5 10 15

Gly Asp Lys Gly Thr Ser Trp Tyr Ile Ile Trp Lys Gly Ser Val Asn 20 25 30

Val Val Thr His Gly Lys Gly Leu Val Thr Thr Leu His Glu Gly Asp 35 40 45

Asp Phe Gly Gln Leu Ala Leu Val Asn Asp Ala Pro Arg Ala Ala Thr 50 55 60

Ile Ile Leu Arg Glu Asp Asn Cys His Phe Leu Arg Val Asp Lys Gln 65 70 75 80

Asp Phe Asn Arg Ile Ile Lys Asp Val Glu Ala Lys Thr Met Arg Leu 85 90 95

Glu Glu His

<210> 23

<211> 97

<212> PRT

<213> Homo sapiens

<400> 23

Ala Met Phe Pro Val Thr His Ile Ala Gly Glu Thr Val Ile Gln Gln 1 5 10 15

Gly Asn Glu Gly Asp Asn Phe Tyr Val Val Asp Gln Gly Glu Val Asp 20 25 30

Val Tyr Val Asn Gly Glu Trp Val Thr Asn Ile Ser Glu Gly Gly Ser 35 40 45

Phe Gly Glu Leu Ala Leu Ile Tyr Gly Thr Pro Arg Ala Ala Thr Val 50 55 60

Lys Ala Lys Thr Asp Leu Lys Leu Trp Gly Ile Asp Arg Asp Ser Tyr 65 70 75 80

Arg Arg Ile Leu Met Gly Ser Thr Leu Arg Lys Arg Lys Met Tyr Glu 85 90 95

Glu

<210> 24

<211> 97

<212> PRT

<213> Homo sapiens

<400> 24

Cys Met Tyr Gly Arg Asn Tyr Gln Gln Gly Ser Tyr Ile Ile Lys Gln
1 5 10 15

Gly Glu Pro Gly Asn His Ile Phe Val Leu Ala Glu Gly Arg Leu Glu
20 25 30

Val Phe Gln Gly Glu Lys Leu Leu Ser Ser Ile Pro Met Trp Thr Thr 35 40 45

Phe Gly Glu Leu Ala Ile Leu Tyr Asn Cys Thr Arg Thr Ala Ser Val 50 55 60

Lys Ala Ile Thr Asn Val Lys Thr Trp Ala Leu Asp Arg Glu Val Phe 65 70 75 80

Gln Asn Ile Met Arg Arg Thr Ala Gln Ala Arg Asp Glu Gln Tyr Arg 85 90 95

Asn

```
<210> 25
```

<211> 103

<212> PRT

<213> Mus musculus

<400> 25

Arg Leu Arg Ser Val Val Tyr Leu Pro Asn Asp Tyr Val Cys Lys
1 5 10 15

Gly Glu Ile Gly Arg Glu Met Tyr Ile Ile Gln Ala Gly Gln Val Gln
20 25 30

Val Leu Gly Gly Pro Asp Gly Lys Ser Val Leu Val Thr Leu Lys Ala 35 40 45

Gly Ser Val Phe Gly Glu Ile Ser Leu Leu Ala Val Gly Gly Gly Asn 50 55 60

Arg Arg Thr Ala Asn Val Val Ala His Gly Phe Thr Asn Leu Phe Ile 65 70 75 80

Leu Asp Lys Lys Asp Leu Asn Glu Ile Leu Val His Tyr Pro Glu Ser 85 90 95

Gln Lys Leu Leu Arg Lys Lys 100

<210> 26

<211> 91

<212> PRT

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: unavailable

<400> 26

Arg Glu Asp Phe Glu Ile Ile Arg Val Phe Asp Gly Asn Asn Ser Tyr 1 5 10 15

Arg Ser Gln Ile Ser Arg Asn Ile Val Val Ala Lys His Val Ser Val 20 25 30

Gln Gln Val Arg Asp Ala Ala Leu Arg Arg Phe His Ile Asn Asp Thr 35 40 45

Pro Glu Arg Tyr Tyr Ile Thr Gln Val Val Glu Glu Glu Glu 50 55 60

Ile Leu Glu Asp Pro Val Pro Leu Arg Asn Val Lys Arg Pro Glu Gly 65 70 75 80

Lys Arg Ala Gln Ile Phe Ile Arg Tyr Tyr Asp

<210> 27

<211> 129

<212> PRT

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: unavailable

<400> 27

Ser Ile Leu Val Thr Ser Gln Asp Lys Ala Pro Ser Val Ile Ser Arg 1 5 10 15

Val Leu Lys Lys Asn Asn Arg Asp Ser Ala Val Ala Ser Glu Tyr Glu 20 25 30

Leu Val Gln Leu Leu Pro Gly Glu Arg Glu Leu Thr Ile Pro Ala Ser 35 40 45

Ala Asn Val Phe Tyr Ala Met Asp Gly Ala Ser His Asp Phe Leu Leu 50 55 60

Arg His Gly Glu Gly Pro Leu Leu Leu His Leu Ala Ser Pro Val Ala 65 70 75 80

Arg Leu Pro Gln Glu Leu Leu Arg Val Arg Glu Glu Gly Ala Pro Phe 85 90 95

Pro Gly Ser Arg Pro Gln Gly Gly Arg Leu His Gly His Cys Ser Glu 100 105 110

Glu Glu Ala Pro Leu Ala Tyr Arg Ser His Gly Val His Thr Arg Cys 115 120 125

Gly